

CRF Error Corrected by the STIC Systems Branch

OIPF 0590
0425
#71

Serial Number: 09/960,632A

CRF Processing Date: 5/1/2002
 Edited by: _____
 Verified by: A (STIC staff)

ENTERED

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: _____
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☒ Deleted: ☒ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☒ Other: corrected global misspellings of primer

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

A-11



RAW SEQUENCE LISTING

DATE: 05/01/2002

PATENT APPLICATION: US/09/960,632A

TIME: 13:36:17

Input Set : A:\Pto.amc

Output Set: N:\CRF3\05012002\I960632A.raw

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3 <110> APPLICANT: PATTOU, Francois
4 KERR-CONTE PATTOU, Julie
6 <120> TITLE OF INVENTION: PROCESS FOR OBTAINING MAMMALIAN INSULIN SECRETING CELLS IN
VITRO AND
7 THEIR USES
9 <130> FILE REFERENCE: B8177-US March 29, 2002
11 <140> CURRENT APPLICATION NUMBER: US 09/960,632A
13 <141> CURRENT FILING DATE: 2001-09-21
15 <150> PRIOR APPLICATION NUMBER: FR00/12547
17 <151> PRIOR FILING DATE: 2000-10-02
19 <160> NUMBER OF SEQ ID NOS: 6
21 <170> SOFTWARE: PatentIn version 3.1
24 <210> SEQ ID NO: 1
25 <211> LENGTH: 18
26 <212> TYPE: DNA
27 <213> ORGANISM: artificial sequence
W--> 28 <220> FEATURE:
29 <223> OTHER INFORMATION: Artificial sequence are primer designed to amplify sequences
W--> 32 <220> FEATURE:
W--> 33 <221> NAME/KEY: primer
34 <222> LOCATION: (1)..(18)
35 <223> OTHER INFORMATION: primer to amplify IPF-1 factor
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39 ccatggatga agtctacc 18
43 <210> SEQ ID NO: 2
44 <211> LENGTH: 19
45 <212> TYPE: DNA
46 <213> ORGANISM: artificial sequence
W--> 47 <220> FEATURE:
48 <223> OTHER INFORMATION: Artificial sequence are primer designed to amplify sequences
W--> 51 <220> FEATURE:
W--> 52 <221> NAME/KEY: primer
53 <222> LOCATION: (1)..(19)
54 <223> OTHER INFORMATION: primer to amplify IPF-1 factor
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62 <210> SEQ ID NO: 3
63 <211> LENGTH: 18
64 <212> TYPE: DNA
65 <213> ORGANISM: artificial sequence
W--> 66 <220> FEATURE:
67 <223> OTHER INFORMATION: Artificial sequence are primer designed to amplify sequences
W--> 70 <220> FEATURE:
W--> 71 <221> NAME/KEY: primer

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RAW SEQUENCE LISTING

DATE: 05/01/2002

PATENT APPLICATION: US/09/960,632A

TIME: 13:36:17

Input Set : A:\Pto.amc

Output Set: N:\CRF3\05012002\I960632A.raw

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72 <222> LOCATION: (1)..(18)
73 <223> OTHER INFORMATION: Primer to amplify insulin
76 <400> SEQUENCE: 3
77 tgtgaaccaa cacctgtg                                     18
81 <210> SEQ ID NO: 4
82 <211> LENGTH: 17
83 <212> TYPE: DNA
84 <213> ORGANISM: artificial sequence
W--> 85 <220> FEATURE:
86 <223> OTHER INFORMATION: Artificial sequence are primer designed to amplify sequences
W--> 89 <220> FEATURE:
W--> 90 <221> NAME/KEY: artificial
91 <222> LOCATION: (1)..(17)
92 <223> OTHER INFORMATION: Primer to amplify insulin
95 <400> SEQUENCE: 4
96 cgtctagttg cagtagt                                     17
100 <210> SEQ ID NO: 5
101 <211> LENGTH: 19
102 <212> TYPE: DNA
103 <213> ORGANISM: artificial sequence
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105 <223> OTHER INFORMATION: Artificial sequence are primer designed to amplify sequences
W--> 108 <220> FEATURE:
W--> 109 <221> NAME/KEY: primer
110 <222> LOCATION: (1)..(19)
111 <223> OTHER INFORMATION: Primer to amplify beta-actin
114 <400> SEQUENCE: 5
115 atcatgtttg agacctcca                                     19
119 <210> SEQ ID NO: 6
120 <211> LENGTH: 20
121 <212> TYPE: DNA
122 <213> ORGANISM: artificial sequence
W--> 123 <220> FEATURE:
124 <223> OTHER INFORMATION: Artificial sequence are primer designed to amplify sequences
W--> 127 <220> FEATURE:
W--> 128 <221> NAME/KEY: primer
129 <222> LOCATION: (1)..(20)
130 <223> OTHER INFORMATION: primer to amplify beta-actin
133 <400> SEQUENCE: 6
134 catctcttgc tcgaagtcca                                     20

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VERIFICATION SUMMARY

DATE: 05/01/2002

PATENT APPLICATION: US/09/960,632A

TIME: 13:36:18

Input Set : A:\Pto.amc

Output Set: N:\CRF3\05012002\I960632A.raw

L:28 M:283 W: Missing Blank Line separator, <220> field identifier
L:33 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1
L:47 M:283 W: Missing Blank Line separator, <220> field identifier
L:52 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:2
L:66 M:283 W: Missing Blank Line separator, <220> field identifier
L:71 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:3
L:85 M:283 W: Missing Blank Line separator, <220> field identifier
L:90 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:4
L:104 M:283 W: Missing Blank Line separator, <220> field identifier
L:109 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:5
L:123 M:283 W: Missing Blank Line separator, <220> field identifier
L:128 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:6



OIPE

RAW SEQUENCE LISTING

DATE: 04/26/2002

PATENT APPLICATION: US/09/960,632A

TIME: 11:54:19

Input Set : A:\PTO.VSK.txt

Output Set: N:\CRF3\04262002\I960632A.raw

Does Not Comply
Corrected Diskette Needed

3 <110> APPLICANT: PATTOU, Francois KERR-CONTE PATTOU, Julie
 5 <120> TITLE OF INVENTION: PROCESS FOR OBTAINING MAMMALIAN INSULIN SECRETING CELLS IN
 VITRO AND
 6 THEIR USES
 8 <130> FILE REFERENCE: B8177-US March 29, 2002
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/09/960,632A
 C--> 12 <141> CURRENT FILING DATE: 2002-04-11
 14 <150> PRIOR APPLICATION NUMBER: FR00/12547
 16 <151> PRIOR FILING DATE: 2000-10-02
 18 <160> NUMBER OF SEQ ID NOS: 6
 20 <170> SOFTWARE: PatentIn version 3.1

ERRORED SEQUENCES

118 <210> SEQ ID NO: 6
 119 <211> LENGTH: 20
 120 <212> TYPE: DNA
 121 <213> ORGANISM: artificial sequence
 W--> 122 <220> FEATURE:
 123 <223> OTHER INFORMATION: Artificial sequence are primer designed to amplify sequences
 W--> 126 <220> FEATURE:
 W--> 127 <221> NAME/KEY: primer
 128 <222> LOCATION: (1)..(20)
 129 <223> OTHER INFORMATION: primer to amplify beta-actin
 132 <400> SEQUENCE: 6
 133 catctcttgc tcgaagtcca 20
 E--> 136 2
 E--> 139 1

VERIFICATION SUMMARY

DATE: 04/26/2002

PATENT APPLICATION: US/09/960,632A

TIME: 11:54:20

Input Set : A:\PTO.VSK.txt

Output Set: N:\CRF3\04262002\I960632A.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application Number
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:27 M:283 W: Missing Blank Line separator, <220> field identifier
L:32 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1
L:46 M:283 W: Missing Blank Line separator, <220> field identifier
L:51 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:2
L:65 M:283 W: Missing Blank Line separator, <220> field identifier
L:70 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:3
L:84 M:283 W: Missing Blank Line separator, <220> field identifier
L:89 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:4
L:103 M:283 W: Missing Blank Line separator, <220> field identifier
L:108 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:5
L:122 M:283 W: Missing Blank Line separator, <220> field identifier
L:127 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:6
L:136 M:254 E: No. of Bases conflict, this line has no nucleotides.
M:254 Repeated in SeqNo=6